

SCORE Search Results Details for Application 10526521 and Search Result 20070808_155054_us-10-526-521a-15.rng.

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This page gives you Search Results detail for the Application 10526521 and Search Result 20070808_155054_us-10-526-521a-15.rng.

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GenCore version 6.2.1
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OM nucleic - nucleic search, using sw model

Run on: August 9, 2007, 18:15:57 ; Search time 357 Seconds
(without alignments)
269.853 Million cell updates/sec

Title: US-10-526-521A-15
Perfect score: 13
Sequence: 1 tggtgagtcacca 13

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5620219 seqs, 3705283702 residues

Total number of hits satisfying chosen parameters: 11240438

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : N_Geneseq_200701:*
1: geneseqn1980s:*
2: geneseqn1990s:*
3: geneseqn2000s:*
4: geneseqn2001as:*
5: geneseqn2001bs:*
6: geneseqn2002as:*
7: geneseqn2002bs:*
8: geneseqn2003as:*
9: geneseqn2003bs:*
10: geneseqn2003cs:*
11: geneseqn2003ds:*
12: geneseqn2004as:*
13: geneseqn2004bs:*
14: geneseqn2005s:*
15: geneseqn2006s:*

16: geneseqn2007s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result			%				DB	ID	Description
	No.	Score	Query Match	Length					
c	1	13	100.0	13	12	ADO57900		Ado57900	Transcrip
	2	13	100.0	13	12	ADO57879		Ado57879	Transcrip
	3	13	100.0	15	12	ADO57904		Ado57904	Transcrip
c	4	13	100.0	15	12	ADO57899		Ado57899	Transcrip
	5	13	100.0	15	12	ADO57880		Ado57880	Transcrip
c	6	13	100.0	15	12	ADO57875		Ado57875	Transcrip
	7	13	100.0	17	12	ADO57903		Ado57903	Transcrip
c	8	13	100.0	17	12	ADO57876		Ado57876	Transcrip
c	9	13	100.0	17	12	ADO57898		Ado57898	Transcrip
	10	13	100.0	17	12	ADO57881		Ado57881	Transcrip
c	11	13	100.0	19	12	ADO57897		Ado57897	Transcrip
	12	13	100.0	19	12	ADO57882		Ado57882	Transcrip
c	13	13	100.0	19	12	ADQ36695		Adq36695	Primer 3
c	14	13	100.0	21	12	ADO57896		Ado57896	Transcrip
	15	13	100.0	21	12	ADO57883		Ado57883	Transcrip
c	16	13	100.0	116	12	ACH89045		Ach89045	Human gen
c	17	13	100.0	231	4	AAL21284		Aal21284	Human bre
	18	13	100.0	250	14	AEH10788		Aeh10788	RAS homol

```

<!--StartFragment-->RESULT 17
AAL21284/c
ID   AAL21284 standard; cDNA; 231 BP.
XX
AC   AAL21284;
XX
DT   07-DEC-2001   (first entry)
XX
DE   Human breast cancer expressed polynucleotide 13741.
XX
KW   Human; breast cancer; cell marker; cytostatic; ss.
XX
OS   Homo sapiens.
XX
PN   WO200151628-A2.
XX
PD   19-JUL-2001.
XX
PF   10-JAN-2001; 2001WO-US000798.
XX
PR   14-JAN-2000; 2000US-0176077P.
PR   14-MAR-2000; 2000US-0189167P.
PR   24-MAR-2000; 2000US-0192099P.
PR   29-MAR-2000; 2000US-0193480P.
PR   15-MAY-2000; 2000US-0205230P.
PR   09-JUN-2000; 2000US-0211315P.
PR   25-JUL-2000; 2000US-0220534P.
XX
PA   (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
XX
PI   Lillie J, Xu Y, Wang Y, Steinmann K;
XX
DR   WPI; 2001-451856/48.
XX
PT   New peptide useful as a marker for the diagnosis of breast cancer.
XX
PS   Claim 1; Page 2444; 3695pp; English.
XX
CC   The invention relates to human breast cancer expressed polynucleotides
CC   (AAL07544-AAL26789) and methods of assessing whether a patient is
CC   afflicted with breast cancer by examining the correlation between the
CC   expression of certain markers and the cancerous state of breast cells.
CC   The polynucleotides and encoded polypeptides are potential markers for
CC   detecting, diagnosing, monitoring, characterising treating and
CC   potentially preventing breast cancer. The polynucleotides and encoded
CC   polypeptides are also useful for isolating compounds with cytostatic
CC   activity
XX
SQ   Sequence 231 BP; 62 A; 67 C; 49 G; 53 T; 0 U; 0 Other;

Query Match          100.0%; Score 13; DB 4; Length 231;
Best Local Similarity 100.0%; Pred. No. 5.9e+02;
Matches 13; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 TGGTGAGTCACCA 13
        |||||
Db      25 TGGTGAGTCACCA 13

```

<!--EndFragment-->